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Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Durreshwar Anjum

Timestamp: Tue Sep 25 12:12:43 EDT 2007

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Reviewer Comments:

? ?

Delete the Non Ascii characters appears above the headings.

?? ?? ?C# hC# h * + O s ? ? ? ? N P ? ?

/

0

>

?

Please delete the end of file text which appears below seq id 656
And save the file in ASCII txt mode

delete the extra heading inserted in the sequence listing which appears
one below the other.

Application No:	10576331	Version No:	1.0
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Input Set:

Output Set:

Started: 2007-09-12 13:26:32.596
Finished: 2007-09-12 13:27:06.289
Elapsed: 0 hr(s) 0 min(s) 33 sec(s) 693 ms
Total Warnings: 2355
Total Errors: 342
No. of SeqIDs Defined: 656
Actual SeqID Count: 656

Error code	Error Description
W 402	Undefined organism found in <213> in SEQ ID (1)
W 402	Undefined organism found in <213> in SEQ ID (2)
W 402	Undefined organism found in <213> in SEQ ID (3)
W 402	Undefined organism found in <213> in SEQ ID (4)
W 402	Undefined organism found in <213> in SEQ ID (5)
W 402	Undefined organism found in <213> in SEQ ID (6)
W 402	Undefined organism found in <213> in SEQ ID (7)
W 402	Undefined organism found in <213> in SEQ ID (8)
W 402	Undefined organism found in <213> in SEQ ID (9)
W 213	Artificial or Unknown found in <213> in SEQ ID (10)
W 213	Artificial or Unknown found in <213> in SEQ ID (11)
W 402	Undefined organism found in <213> in SEQ ID (12)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (12)
W 402	Undefined organism found in <213> in SEQ ID (13)
W 213	Artificial or Unknown found in <213> in SEQ ID (14)
W 213	Artificial or Unknown found in <213> in SEQ ID (15)
W 213	Artificial or Unknown found in <213> in SEQ ID (16)
W 213	Artificial or Unknown found in <213> in SEQ ID (17)
W 402	Undefined organism found in <213> in SEQ ID (18)

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Error code	Error Description
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (34)
W 402	Undefined organism found in <213> in SEQ ID (37)
W 402	Undefined organism found in <213> in SEQ ID (38)
W 402	Undefined organism found in <213> in SEQ ID (40)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (40)
W 213	Artificial or Unknown found in <213> in SEQ ID (45)
W 213	Artificial or Unknown found in <213> in SEQ ID (46)
W 213	Artificial or Unknown found in <213> in SEQ ID (47)
W 213	Artificial or Unknown found in <213> in SEQ ID (48)
W 213	Artificial or Unknown found in <213> in SEQ ID (49)
W 213	Artificial or Unknown found in <213> in SEQ ID (50)
W 213	Artificial or Unknown found in <213> in SEQ ID (51)
W 213	Artificial or Unknown found in <213> in SEQ ID (52)
W 402	Undefined organism found in <213> in SEQ ID (61)
W 402	Undefined organism found in <213> in SEQ ID (62)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (62)
W 402	Undefined organism found in <213> in SEQ ID (65)
W 402	Undefined organism found in <213> in SEQ ID (66)
W 402	Undefined organism found in <213> in SEQ ID (69) This error has occurred more than 20 times, will not be displayed

Input-Set:**Output Set:**

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Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (79)
W 213	Artificial or Unknown found in <213> in SEQ ID (80)
W 213	Artificial or Unknown found in <213> in SEQ ID (81)
W 213	Artificial or Unknown found in <213> in SEQ ID (82)
W 213	Artificial or Unknown found in <213> in SEQ ID (83)
W 213	Artificial or Unknown found in <213> in SEQ ID (84) This error has occurred more than 20 times, will not be displayed
E 341	'Xaa' position not defined SEQID (125) POS (170)
E 341	'Xaa' position not defined SEQID (125) POS (171)
E 341	'Xaa' position not defined SEQID (125) POS (172)
E 341	'Xaa' position not defined SEQID (125) POS (173)
E 341	'Xaa' position not defined SEQID (125) POS (174)
E 341	'Xaa' position not defined SEQID (125) POS (175)
E 341	'Xaa' position not defined SEQID (125) POS (176)
E 341	'Xaa' position not defined SEQID (125) POS (177)
E 341	'Xaa' position not defined SEQID (125) POS (178)
E 341	'Xaa' position not defined SEQID (125) POS (179)
E 341	'Xaa' position not defined SEQID (125) POS (180)
E 341	'Xaa' position not defined SEQID (125) POS (181)
E 341	'Xaa' position not defined SEQID (125) POS (182)
E 341	'Xaa' position not defined SEQID (125) POS (183)
E 341	'Xaa' position not defined SEQID (125) POS (184)
E 341	'Xaa' position not defined SEQID (125) POS (185)

Input Set:

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Error code	Error Description
E 341	'Xaa' position not defined SEQID (125) POS (186)
E 341	'Xaa' position not defined SEQID (125) POS (187)
E 341	'Xaa' position not defined SEQID (125) POS (188)
E 341	'Xaa' position not defined SEQID (125) POS (189) This error has occurred more than 20 times, will not be displayed
W 112	Upper case found in data; Found at position(977) SEQID(656)
W 112	Upper case found in data; Found at position(1528) SEQID(656)
W 112	Upper case found in data; Found at position(1930) SEQID(656)
W 112	Upper case found in data; Found at position(1950) SEQID(656)
W 112	Upper case found in data; Found at position(2002) SEQID(656)
W 112	Upper case found in data; Found at position(2030) SEQID(656)
W 112	Upper case found in data; Found at position(2440) SEQID(656)
W 112	Upper case found in data; Found at position(2452) SEQID(656)
W 112	Upper case found in data; Found at position(2496) SEQID(656)
W 112	Upper case found in data; Found at position(2500) SEQID(656)
W 112	Upper case found in data; Found at position(2504) SEQID(656)
W 112	Upper case found in data; Found at position(2531) SEQID(656)
W 112	Upper case found in data; Found at position(2552) SEQID(656)
W 112	Upper case found in data; Found at position(2941) SEQID(656)
W 112	Upper case found in data; Found at position(2945) SEQID(656)
W 112	Upper case found in data; Found at position(2966) SEQID(656)
W 112	Upper case found in data; Found at position(2991) SEQID(656)
W 112	Upper case found in data; Found at position(3015) SEQID(656)

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Error code	Error Description
W 112	Upper case found in data; Found at position(3043) SEQID(656)
W 112	Upper case found in data; Found at position(3047) SEQID(656) This error has occurred more than 20 times, will not be displayed
E 342	'n' position not defined found at POS: 4496 SEQID(656)
E 342	'n' position not defined found at POS: 9429 SEQID(656)
E 342	'n' position not defined found at POS: 12391 SEQID(656)
E 342	'n' position not defined found at POS: 12931 SEQID(656)
E 342	'n' position not defined found at POS: 19721 SEQID(656)
E 342	'n' position not defined found at POS: 30772 SEQID(656)
E 342	'n' position not defined found at POS: 30776 SEQID(656)
E 342	'n' position not defined found at POS: 30779 SEQID(656)
E 342	'n' position not defined found at POS: 30783 SEQID(656)
E 342	'n' position not defined found at POS: 30787 SEQID(656)
E 342	'n' position not defined found at POS: 30791 SEQID(656)
E 342	'n' position not defined found at POS: 30795 SEQID(656)
E 342	'n' position not defined found at POS: 30799 SEQID(656)
E 342	'n' position not defined found at POS: 30803 SEQID(656)
E 342	'n' position not defined found at POS: 30807 SEQID(656)
E 342	'n' position not defined found at POS: 30811 SEQID(656)
E 342	'n' position not defined found at POS: 30815 SEQID(656)
E 342	'n' position not defined found at POS: 30819 SEQID(656)
E 342	'n' position not defined found at POS: 30823 SEQID(656)
E 342	'n' position not defined found at POS: 30827 SEQID(656)

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Error code	Error Description
E 259	Found undefined lettercode; POS (90993) SEQID(656)
E 259	Found undefined lettercode; POS (90994) SEQID(656)
E 259	Found undefined lettercode; POS (90995) SEQID(656)
E 259	Found undefined lettercode; POS (90996) SEQID(656)
E 259	Found undefined lettercode; POS (90997) SEQID(656)
E 259	Found undefined lettercode; POS (90998) SEQID(656)
E 259	Found undefined lettercode; POS (90999) SEQID(656)
E 259	Found undefined lettercode; POS (91000) SEQID(656)
E 259	Found undefined lettercode; POS (91001) SEQID(656)
E 259	Found undefined lettercode; POS (91002) SEQID(656)
E 259	Found undefined lettercode; POS (91003) SEQID(656)
E 259	Found undefined lettercode; POS (91004) SEQID(656)
E 259	Found undefined lettercode; POS (91005) SEQID(656)
E 259	Found undefined lettercode; POS (91006) SEQID(656)
E 259	Found undefined lettercode; POS (91008) SEQID(656)
E 259	Found undefined lettercode; POS (91009) SEQID(656)
E 259	Found undefined lettercode; POS (91010) SEQID(656)
E 259	Found undefined lettercode; POS (91011) SEQID(656)
E 259	Found undefined lettercode; POS (91012) SEQID(656)
E 259	Found undefined lettercode; POS (91013) SEQID(656)

This error has occurred more than 20 times, will not be displayed

<210> 1

<211> 1680

<212> DNA

<213> Cellulomonas strain 69B4

<400> 1

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accagcacgc tcctaacctc cgagcaccgg cgaccgccgg gtgcgatgaa agggacgaac 180
cgagatgaca ccacgcacag tcacgcgggc cctggccgtg gccaccgcag ccgccacact 240
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accgccacgc ctggccgaga agctcgaccc cgacctctc gagggcatgg agcgcgacct 360
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cggcgaggcc ctgcgcgaag agctcgacga ggacttcgcc ggcacctggg tcgaggacga 480
cgtcctgtac gtcgccacca ccgacgagga cgccgtcgag gaggtcgagg gcgaaggcgc 540
cacggccgtc accgtcgagc actccctggc cgacctcgag gcctggaaga ccgtcctcga 600
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cgtcgtcgtc gccgtcaagg ccggagccca ggacgtcgcc gccggcctcg tcgaaggtgc 720
cgacgtcccg tccgacgcgg tgaccttctg cgagaccgac gagaccccgc ggacctggt 780
cgacgtgatc ggcggaacg cctacaccat cggggggcgc agccgctgct cgatcggggt 840
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gctcgccggc aaccaggccc agggcgctac gtcggcgggc tccggcaact gccgcaccgg 1260
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caccacggac tcgggcagca gcccgcccc tgcaccgacc tcttgaccg gctacgccc 1380
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ccccggtcc aacgagacca tcacctacc cggaacgcc ggctactacc gctacgtggt 1620
caacgcgcgc tccggctccg gtgcctacac catggggctc accctcccct gacgtagcgc 1680
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<210> 2

<211> 1488

<212> DNA

<213> Cellulomonas strain 69B4

<400> 2

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ccacgcctgg ccgagaagct cgaccccgac ctctcgagg ccatggagcg cgacctgggc 180
ctcgagcggg aggaagcgc cgccacctg gcgttcacg acgacgcagc cgagaccggc 240
gaggccctcg ccgaagagct cgacgaggac ttgcgggca cctgggtcga ggacgacgtc 300
ctgtacgtcg ccaccaccga cgaggacgcc gtcgaggagg tcgagggcga aggcgccacg 360
gccgtcaccg tcgagcactc cctggccgac ctcgaggcct ggaagaccgt cctcgacgcc 420
gccctcgagg gccacgacga cgtgcccacc tggtagctcg acgtcccgc caacagcgtc 480
gtcgtcgccg tcaaggccgg agcccaggac gtcgcccgcc gcctcgtcga aggtgccgac 540
gtcccgctcc acgcccgtgac cttcgtcgag accgacgaga ccccgcgga catgttcgac 600
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gtcaacggcg ggttcacac cgccggccac tgcggcgca ccggcgccac caccgccaac 720
cccaccggga ccttcgcccg gtccagcttc ccgggcaacg actacgcgtt cgtccgtacc 780
ggggccggcg tgaacctgct ggcccaggtc aacaactact ccggtggccg cgtccaggtc 840
gccgggcaca ccgcggcccc cgtcggtcgt gccgtgtgcc ggtccgggtc gaccaccggg 900
tggcactgcg gcaccatcac tgcgtcaac tctcgggtca cctacccga gggcaccgtc 960
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accacgttct	tccagccggt	caaccccatc	ctccaggcgt	acggcctgag	gatgatcacc	1140
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gtcaaccggt	ccgggaccca	cagcgtgtgc	ctcaacgggc	cctccggtgc	ggacttcgac	1320
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ggctccaacg	agaccatcac	ctaccgcggc	aacgcggct	actaccgcta	cgtggtcaac	1440
gcccggtccg	gctccggtgc	ctacaccatg	gggtcacc	tcccctga		1488

<210> 3
 <211> 1404
 <212> DNA
 <213> Cellulomonas spp.

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accctggcgt	tccagcacga
gaggacttcg	ccggcacctg
gacgccgtcg	aggaggtcga
gccgacctcg	aggcctggaa
cccacctggt	acgtcgacgt
caggacgtcg	ccgccggcct
gtcgagaccg	acgagacccc
atcggggggc	gcagccgctg
ggccactgcg	gccgcaccgg
agcttcccgg	gcaacgacta
caggtaacaa	actactccgg
ggctcgcccg	tgtgccggtc
ctcaactcct	cggtcaccta
tgcgccgagc	ccggcgactc
acgtccggcg	gctccggcaa
cccatcctcc	aggcgtacgg
cctgcaccga	cctcctgcac
cgggccgccc	cccagcccaa
gtgtgcctca	acgggcccctc
agctcctggg	tgaccgtcgc
cgcggaacg	ccggctaacta
accatggggc	tcacctccc
	ctga
	60
	120
	180
	240
	300
	360
	420
	480
	540
	600
	660
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	960
	1020
	1080
	1140
	1200
	1260
	1320
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	1404

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 <211> 567
 <212> DNA
 <213> Cellulomonas spp.

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accgggtggc	actgcgga
accgtccggc	gctgatccg
ctgctcgccc	gcaaccaggc
ggtggcacca	cggttctcca
atcaccacgg	actcgggcag
	cagcccggcc
	cgcgcccgcc
	gaccacagc
	ctggaacggc
	catcacctac
	cgggtgcctac
	1404
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	120
	180
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	300
	360
	420
	480
	540
	567

<210> 5
<211> 83
<212> DNA
<213> Cellulomonas strain 69B4

<400> 5
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60
83

<210> 6
<211> 495
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<213> Cellulomonas strain 69B4

<400> 6

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20 25 30
Pro Pro Gly Ser Ala Ser Ala Pro Pro Arg Leu Ala Glu Lys Leu Asp
35 40 45
Pro Asp Leu Leu Glu Ala Met Glu Arg Asp Leu Gly Leu Asp Ala Glu
50 55 60
Glu Ala Ala Ala Thr Leu Ala Phe Gln His Asp Ala Ala Glu Thr Gly
65 70 75 80
Glu Ala Leu Ala Glu Glu Leu Asp Glu Asp Phe Ala Gly Thr Trp Val
85 90 95
Glu Asp Asp Val Leu Tyr Val Ala Thr Thr Asp Glu Asp Ala Val Glu
100 105 110
Glu Val Glu Gly Glu Gly Ala Thr Ala Val Thr Val Glu His Ser Leu
115 120 125
Ala Asp Leu Glu Ala Trp Lys Thr Val Leu Asp Ala Ala Leu Glu Gly
130 135 140
His Asp Asp Val Pro Thr Trp Tyr Val Asp Val Pro Thr Asn Ser Val
145 150 155 160
Val Val Ala Val Lys Ala Gly Ala Gln Asp Val Ala Ala Gly Leu Val
165 170 175
Glu Gly Ala Asp Val Pro Ser Asp Ala Val Thr Phe Val Glu Thr Asp
180 185 190
Glu Thr Pro Arg Thr Met Phe Asp Val Ile Gly Gly Asn Ala Tyr Thr
195 200 205
Ile Gly Gly Arg Ser Arg Cys Ser Ile Gly Phe Ala Val Asn Gly Gly
210 215 220
Phe Ile Thr Ala Gly His Cys Gly Arg Thr Gly Ala Thr Thr Ala Asn
225 230 235 240
Pro Thr Gly Thr Phe Ala Gly Ser Ser Phe Pro Gly Asn Asp Tyr Ala
245 250 255
Phe Val Arg Thr Gly Ala Gly Val Asn Leu Leu Ala Gln Val Asn Asn
260 265 270
Tyr Ser Gly Gly Arg Val Gln Val Ala Gly His Thr Ala Ala Pro Val
275 280 285
Gly Ser Ala Val Cys Arg Ser Gly Ser Thr Thr Gly Trp His Cys Gly

290

295

300

Thr Ile Thr Ala Leu Asn Ser Ser Val Thr Tyr Pro Glu Gly Thr Val
 305 310 315 320
 Arg Gly Leu Ile Arg Thr Thr Val Cys Ala Glu Pro Gly Asp Ser Gly
 325 330 335
 Gly Ser Leu Leu Ala Gly Asn Gln Ala Gln Gly Val Thr Ser Gly Gly
 340 345 350
 Ser Gly Asn Cys Arg Thr Gly Gly Thr Thr Phe Phe Gln Pro Val Asn
 355 360 365
 Pro Ile Leu Gln Ala Tyr Gly Leu Arg Met Ile Thr Thr Asp Ser Gly
 370 375 380
 Ser Ser Pro Ala Pro Ala Pro Thr Ser Cys Thr Gly Tyr Ala Arg Thr
 385 390 395 400
 Phe Thr Gly Thr Leu Ala Ala Gly Arg Ala Ala Ala Gln Pro Asn Gly
 405 410 415
 Ser Tyr Val Gln Val Asn Arg Ser Gly Thr His Ser Val Cys Leu Asn
 420 425 430
 Gly Pro Ser Gly Ala Asp Phe Asp Leu Tyr Val Gln Arg Trp Asn Gly
 435 440 445
 Ser Ser Trp Val Thr Val Ala Gln Ser Thr Ser Pro Gly Ser Asn Glu
 450 455 460
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 465 470 475 480
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 485 490 495

<210> 7

<211> 467

<212> PRT

<213> Cellulomonas strain 69B4

<400> 7

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 20 25 30
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 35 40 45
 Ala Glu Thr Gly Glu Ala Leu Ala Glu Glu Leu Asp Glu Asp Phe Ala
 50 55 60
 Gly Thr Trp Val Glu Asp Asp Val Leu Tyr Val Ala Thr Thr Asp Glu
 65 70 75 80
 Asp Ala Val Glu Glu Val Glu Gly Glu Gly Ala Thr Ala Val Thr Val
 85 90 95
 Glu His Ser Leu Ala Asp Leu Glu Ala Trp Lys Thr Val Leu Asp Ala
 100 105 110
 Ala Leu Glu Gly His Asp Asp Val Pro Thr Trp Tyr Val Asp Val Pro
 115 120 125
 Thr Asn Ser Val Val Val Ala Val Lys Ala Gly Ala Gln Asp Val Ala
 130 135 140
 Ala Gly Leu Val Glu Gly Ala Asp Val Pro Ser Asp Ala Val Thr Phe
 145 150 155 160
 Val Glu Thr Asp Glu Thr Pro Arg Thr Met Phe Asp Val Ile Gly Gly
 165 170 175
 Asn Ala Tyr Thr Ile Gly Gly Arg Ser Arg Cys Ser Ile Gly Phe Ala
 180 185 190

Val	Asn	Gly	Gly	Phe	Ile	Thr	Ala	Gly	His	Cys	Gly	Arg	Thr	Gly	Ala
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210						215					220				
Asn	Asp	Tyr	Ala	Phe	Val	Arg	Thr	Gly	Ala	Gly	Val	Asn	Leu	Leu	Ala
225					230					235					240
Gln	Val	Asn	Asn	Tyr	Ser	Gly	Gly	Arg	Val	Gln	Val	Ala	Gly	His	Thr
			245					250						255	
Ala	Ala	Pro	Val	Gly	Ser	Ala	Val	Cys	Arg	Ser	Gly	Ser	Thr	Thr	Gly
		260						265					270		
Trp	His	Cys	Gly	Thr	Ile	Thr	Ala	Leu	Asn	Ser	Ser	Val	Thr	Tyr	Pro
	275					280						285			
Glu	Gly	Thr	Val	Arg	Gly	Leu	Ile	Arg	Thr	Thr	Val	Cys	Ala	Glu	Pro
290						295					300				
Gly	Asp	Ser	Gly	Gly	Ser	Leu	Leu	Ala	Gly	Asn	Gln	Ala	Gln	Gly	Val
305					310					315					320
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			325					330						335	
Gln	Pro	Val	Asn	Pro	Ile	Leu	Gln	Ala	Tyr	Gly	Leu	Arg	Met	Ile	Thr
	340							345					350		
Thr	Asp	Ser	Gly	Ser	Ser	Pro	Ala	Pro	Ala	Pro	Thr	Ser	Cys	Thr	Gly
	355					360						365			
Tyr	Ala	Arg	Thr	Phe	Thr	Gly	Thr	Leu	Ala	Ala	Gly	Arg	Ala	Ala	Ala
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Gln	Pro	Asn	Gly	Ser	Tyr	Val	Gln	Val	Asn	Arg	Ser	Gly	Thr	His	Ser
385					390					395					400
Val	Cys	Leu	Asn	Gly	Pro	Ser	Gly	Ala	Asp	Phe	Asp	Leu	Tyr	Val	Gln
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Arg	Trp	Asn	Gly	Ser	Ser	Trp	Val	Thr	Val	Ala	Gln	Ser	Thr	Ser	Pro
		420						425					430		
Gly	Ser	Asn	Glu	Thr	Ile	Thr	Tyr	Arg	Gly	Asn	Ala	Gly	Tyr	Tyr	Arg
	435					440						445			
Tyr	Val	Val	Asn	Ala	Ala	Ser	Gly	Ser	Gly	Ala	Tyr	Thr	Met	Gly	Leu
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Cys	Gly	Arg	Thr	Gly	Ala	Thr	Thr	Ala	Asn	Pro	Thr	Gly	Thr	Phe	Ala
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Gly	Val	Asn	Leu	Leu	Ala	Gln	Val	Asn	Asn	Tyr	Ser	Gly	Gly	Arg	Val
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Gln	Val	Ala	Gly	His	Thr	Ala	Ala	Pro	Val	Gly	Ser	Ala	Val	Cys	Arg
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	100		105		110										
Ser	Ser	Val	Thr	Tyr	Pro	Glu	Gly	Thr	Val	Arg	Gly	Leu	Ile	Arg	Thr
	115						120					125			
Thr	Val	Cys	Ala	Glu	Pro	Gly	Asp	Ser	Gly	Gly	Ser	Leu	Leu	Ala	Gly
	130					135					140				
Asn	Gln	Ala	Gln	Gly	Val	Thr	Ser	Gly	Gly	Ser	Gly	Asn	Cys	Arg	Thr
145					150					155				160	
Gly	Gly	Thr	Thr	Phe	Phe	Gln	Pro	Val	Asn	Pro	Ile	Leu	Gln	Ala	Tyr
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Gly	Leu	Arg	Met	Ile	Thr	Thr	Asp	Ser	Gly	Ser	Ser	Pro			
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Met	Thr	Pro	Arg	Thr	Val	Thr	Arg	Ala	Leu	Ala	Val	Ala	Thr	Ala	Ala
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Asp Gly Trp Asp Cys Gly Thr Ile Thr Ala Leu Asn Ser Ser Val Thr
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Tyr